

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/618,896

Source: IFWO

Date Processed by STIC: 7/14/06

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IFWO

RAW SEQUENCE LISTING

DATE: 07/14/2006

PATENT APPLICATION: US/10/618,896

TIME: 08:34:18

Input Set : A:\960296.00096.txt

Output Set: N:\CRF4\07142006\J618896.raw

3 <110> APPLICANT: Ahlquist, Paul
 4 Ishikawa, Masayuki
 5 Barcelona, Juana
 6 Price, Duane
 7 Lee, Wai-Ming
 9 <120> TITLE OF INVENTION: Yeast genes that affect viral replication
 11 <130> FILE REFERENCE: 960296.00096
 C--> 13 <140> CURRENT APPLICATION NUMBER: US/10/618,896
 C--> 13 <141> CURRENT FILING DATE: 2003-07-14
 13 <160> NUMBER OF SEQ ID NOS: 22
 15 <170> SOFTWARE: PatentIn version 3.3
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 32
 19 <212> TYPE: PRT
 20 <213> ORGANISM: Saccharomyces cerevisiae
 22 <400> SEQUENCE: 1
 24 Leu Arg Val Leu Thr Gln Asp Gly Arg Val Tyr Ile Gly Gln Leu Met
 25 1 5 10 15
 28 Ala Phe Asp Lys His Met Asn Leu Val Leu Asn Glu Cys Ile Glu Glu
 29 20 25 30
 32 <210> SEQ ID NO: 2
 33 <211> LENGTH: 14
 34 <212> TYPE: PRT
 35 <213> ORGANISM: Saccharomyces cerevisiae
 37 <400> SEQUENCE: 2
 39 Leu Gly Leu Thr Ile Leu Arg Gly Glu Gln Ile Leu Ser Thr
 40 1 5 10
 43 <210> SEQ ID NO: 3
 44 <211> LENGTH: 32
 45 <212> TYPE: PRT
 46 <213> ORGANISM: Saccharomyces cerevisiae
 48 <400> SEQUENCE: 3
 50 Val Thr Ile Glu Leu Lys Asn Gly Thr Thr Val Trp Gly Thr Leu Gln
 51 1 5 10 15
 54 Ser Val Ser Pro Gln Met Asn Ala Ile Leu Thr Asp Val Lys Leu Thr
 55 20 25 30
 58 <210> SEQ ID NO: 4
 59 <211> LENGTH: 14
 60 <212> TYPE: PRT
 61 <213> ORGANISM: Saccharomyces cerevisiae
 63 <400> SEQUENCE: 4
 65 Leu Gln Tyr Ile Asn Ile Arg Gly Asn Thr Ile Arg Gln Ile
 66 1 5 10

p.6

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69 <210> SEQ ID NO: 5
70 <211> LENGTH: 32
71 <212> TYPE: PRT
72 <213> ORGANISM: Saccharomyces cerevisiae
74 <400> SEQUENCE: 5
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80 Gly Phe Asp Glu Phe Met Asn Val Val Ile Asp Glu Ala Val Glu Ile
81          20          25          30
84 <210> SEQ ID NO: 6
85 <211> LENGTH: 14
86 <212> TYPE: PRT
87 <213> ORGANISM: Saccharomyces cerevisiae
89 <400> SEQUENCE: 6
91 Leu Gly Lys Ile Leu Leu Lys Gly Asp Asn Ile Thr Leu Ile
92 1          5          10
95 <210> SEQ ID NO: 7
96 <211> LENGTH: 33
97 <212> TYPE: PRT
98 <213> ORGANISM: Saccharomyces cerevisiae
100 <400> SEQUENCE: 7
102 Val Gly Val Lys Leu Lys Phe Asn Ser Thr Glu Tyr Arg Gly Thr Leu
103 1          5          10          15
106 Val Ser Thr Asp Asn Tyr Phe Asn Leu Gln Leu Asn Glu Ala Glu Glu
107          20          25          30
110 Phe
114 <210> SEQ ID NO: 8
115 <211> LENGTH: 14
116 <212> TYPE: PRT
117 <213> ORGANISM: Saccharomyces cerevisiae
119 <400> SEQUENCE: 8
121 Leu Gly Glu Ile Phe Ile Arg Cys Asn Asn Val Leu Tyr Ile
122 1          5          10
125 <210> SEQ ID NO: 9
126 <211> LENGTH: 32
127 <212> TYPE: PRT
128 <213> ORGANISM: Saccharomyces cerevisiae
130 <400> SEQUENCE: 9
132 Ile Leu Leu Asn Ile Asn Gly Ser Arg Lys Val Ala Gly Ile Leu Arg
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137          20          25          30
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141 <211> LENGTH: 14
142 <212> TYPE: PRT
143 <213> ORGANISM: Saccharomyces cerevisiae
145 <400> SEQUENCE: 10
147 Ile Gly Met Val Val Ile Arg Gly Asn Ser Ile Ile Met Leu
148 1          5          10

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151 <210> SEQ ID NO: 11
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153 <212> TYPE: PRT
154 <213> ORGANISM: Saccharomyces cerevisiae
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162 Thr Phe Asp Gln Tyr Ala Asn Leu Ile Leu Gln Asp Cys Val Glu Arg
163 20 25 30
166 <210> SEQ ID NO: 12
167 <211> LENGTH: 14
168 <212> TYPE: PRT
169 <213> ORGANISM: Saccharomyces cerevisiae
171 <400> SEQUENCE: 12
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174 1 5 10
177 <210> SEQ ID NO: 13
178 <211> LENGTH: 31
179 <212> TYPE: PRT
180 <213> ORGANISM: Schizosaccharomyces pombe
182 <400> SEQUENCE: 13
184 Ile Val Val Leu Arg Asp Gly Lys Lys Leu Ile Gly Ile Leu Arg Ser
185 1 5 10 15
188 Phe Asp Gln Phe Ala Asn Leu Met Leu Gln Tyr Thr Ile Glu Arg
189 20 25 30
192 <210> SEQ ID NO: 14
193 <211> LENGTH: 14
194 <212> TYPE: PRT
195 <213> ORGANISM: Schizosaccharomyces pombe
197 <400> SEQUENCE: 14
199 Arg Gly Val Tyr Ile Val Arg Gly Glu Asn Val Val Leu Leu
200 1 5 10
203 <210> SEQ ID NO: 15
204 <211> LENGTH: 31
205 <212> TYPE: PRT
206 <213> ORGANISM: Homo sapiens
208 <400> SEQUENCE: 15
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214 Ile Asp Gln Phe Ala Asn Leu Val Leu His Gln Thr Val Glu Arg
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218 <210> SEQ ID NO: 16
219 <211> LENGTH: 14
220 <212> TYPE: PRT
221 <213> ORGANISM: Homo sapiens
223 <400> SEQUENCE: 16
225 Arg Gly Ile Phe Val Val Arg Gly Glu Asn Val Val Leu Leu
226 1 5 10
229 <210> SEQ ID NO: 17

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230 <211> LENGTH: 31
231 <212> TYPE: PRT
232 <213> ORGANISM: Caenorhabditis elegans
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236 Leu Val Val Leu Arg Asp Gly Arg Lys Leu Ile Gly Phe Leu Arg Ser
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241          20          25          30
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245 <211> LENGTH: 14
246 <212> TYPE: PRT
247 <213> ORGANISM: Caenorhabditis elegans
249 <400> SEQUENCE: 18
251 Gln Gly Phe Met Leu Ile Arg Gly Glu Asn Val Glu Leu Ala
252 1          5          10
255 <210> SEQ ID NO: 19
256 <211> LENGTH: 32
257 <212> TYPE: PRT
258 <213> ORGANISM: Saccharomyces cerevisiae
260 <400> SEQUENCE: 19
262 Leu Ile Val Ser Thr Leu Glu Asp Arg Ile Leu Val Gly Ser Leu Val
263 1          5          10          15
266 Ala Val Asp Ala Gln Met Asn Leu Leu Leu Asp His Val Glu Glu Arg
267          20          25          30
270 <210> SEQ ID NO: 20
271 <211> LENGTH: 14
272 <212> TYPE: PRT
273 <213> ORGANISM: Saccharomyces cerevisiae
275 <400> SEQUENCE: 20
277 Gly Leu Val Ser Val Pro Arg Arg Ser Val Lys Thr Ile Met
278 1          5          10
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282 <211> LENGTH: 33
283 <212> TYPE: PRT
284 <213> ORGANISM: Artificial
286 <220> FEATURE:
287 <223> OTHER INFORMATION: conserved sequence of Sm motif 1
290 <220> FEATURE:
291 <221> NAME/KEY: MISC_FEATURE
292 <222> LOCATION: (1)..(1)
293 <223> OTHER INFORMATION: X is a hydrophobic amino acid
295 <220> FEATURE:
296 <221> NAME/KEY: MISC_FEATURE
297 <222> LOCATION: (2)..(2)
298 <223> OTHER INFORMATION: X can be any amino acid
300 <220> FEATURE:
301 <221> NAME/KEY: MISC_FEATURE
302 <222> LOCATION: (3)..(3)
303 <223> OTHER INFORMATION: X is a hydrophobic amino acid

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305 <220> FEATURE:
306 <221> NAME/KEY: MISC_FEATURE
307 <222> LOCATION: (4)..(4)
308 <223> OTHER INFORMATION: X can be any amino acid
310 <220> FEATURE:
311 <221> NAME/KEY: MISC_FEATURE
312 <222> LOCATION: (6)..(11)
313 <223> OTHER INFORMATION: X can be any amino acid
315 <220> FEATURE:
316 <221> NAME/KEY: MISC_FEATURE
317 <222> LOCATION: (12)..(12)
318 <223> OTHER INFORMATION: X is a hydrophobic amino acid
320 <220> FEATURE:
321 <221> NAME/KEY: MISC_FEATURE
322 <222> LOCATION: (13)..(13)
323 <223> OTHER INFORMATION: X can be any amino acid
325 <220> FEATURE:
326 <221> NAME/KEY: MISC_FEATURE
327 <222> LOCATION: (15)..(15)
328 <223> OTHER INFORMATION: X can be any amino acid
330 <220> FEATURE:
331 <221> NAME/KEY: MISC_FEATURE
332 <222> LOCATION: (16)..(16)
333 <223> OTHER INFORMATION: X is a hydrophobic amino acid
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336 <221> NAME/KEY: MISC_FEATURE
337 <222> LOCATION: (17)..(19)
338 <223> OTHER INFORMATION: X can be any amino acid
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341 <221> NAME/KEY: MISC_FEATURE
342 <222> LOCATION: (21)..(22)
343 <223> OTHER INFORMATION: X can be any amino acid
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346 <221> NAME/KEY: MISC_FEATURE
347 <222> LOCATION: (25)..(25)
348 <223> OTHER INFORMATION: X is a hydrophobic amino acid
350 <220> FEATURE:
351 <221> NAME/KEY: MISC_FEATURE
352 <222> LOCATION: (26)..(26)
353 <223> OTHER INFORMATION: X can be any amino acid
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356 <221> NAME/KEY: MISC_FEATURE
357 <222> LOCATION: (27)..(27)
358 <223> OTHER INFORMATION: X is a hydrophobic amino acid
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361 <221> NAME/KEY: MISC_FEATURE
362 <222> LOCATION: (28)..(31)
363 <223> OTHER INFORMATION: X can be any amino acid
365 <220> FEATURE:

RAW SEQUENCE LISTING ERROR SUMMARY
 PATENT APPLICATION: US/10/618,896

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:21; Xaa Pos. ~~1, 2, 3, 4, 6, 7, 8, 9, 10, 11, 12, 13, 15, 16, 17, 18, 19, 21, 22, 25, 26, 27~~
 Seq#:21; Xaa Pos. ~~28, 29, 30, 31, 33~~
 Seq#:22; Xaa Pos. 1, 3, 4, 5, 6, 9, 11, 12, 13, 14

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete, per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:21,22

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L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:372 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0

M:341 Repeated in SeqNo=21

L:435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0